

FIG.1

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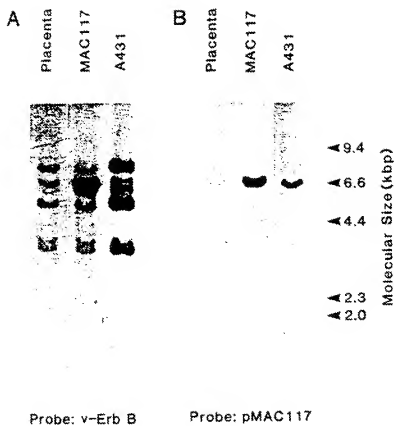


FIG.2

HOMOLOGY

GMSYLEVDRLVHRDLAARNVLKSPNHHVKITDFGLARLLDI
 * * * * * * *
 N R T Q K GA
 N ER I T Q K GA
 V RMNY R A I GENLVG VA I---
 A E KKNI C GENHL VA MTG
 AF AKNCI LT GRVA G DI-M
 A INAKKE CW AHDT F MS DI---

Figure 6. Amino acid sequence alignment of the deduced amino acid sequences of the three proteins. The amino acid residues are numbered from the N-terminus. The asterisks indicate identical amino acids. The arrow indicates the position of the start codon.

FILE 3

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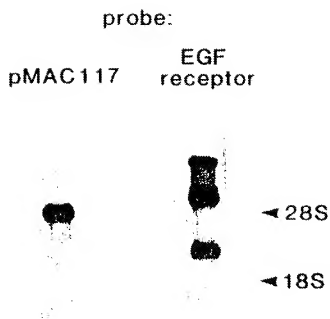
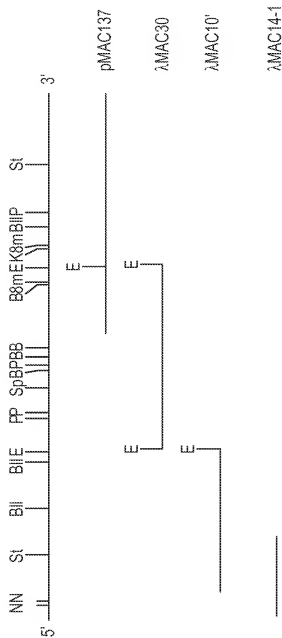


FIG.4

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A



B



FIG.5

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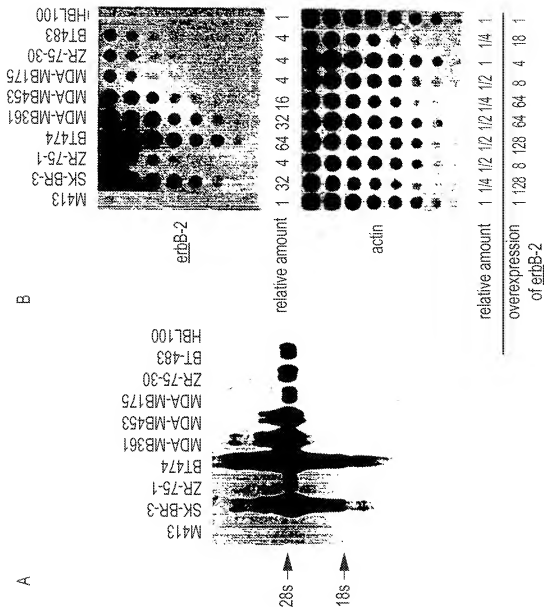


FIG.6



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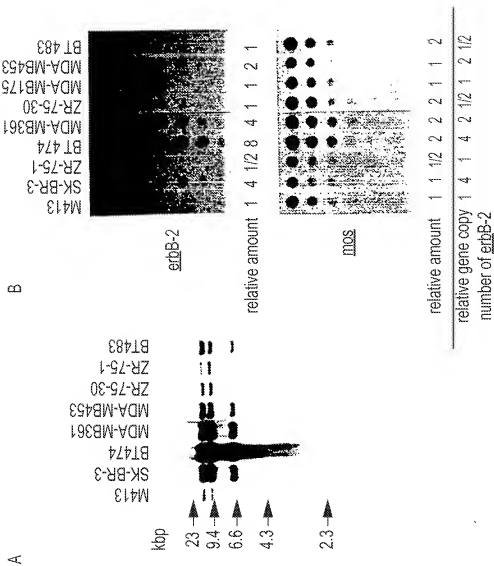


FIG.8

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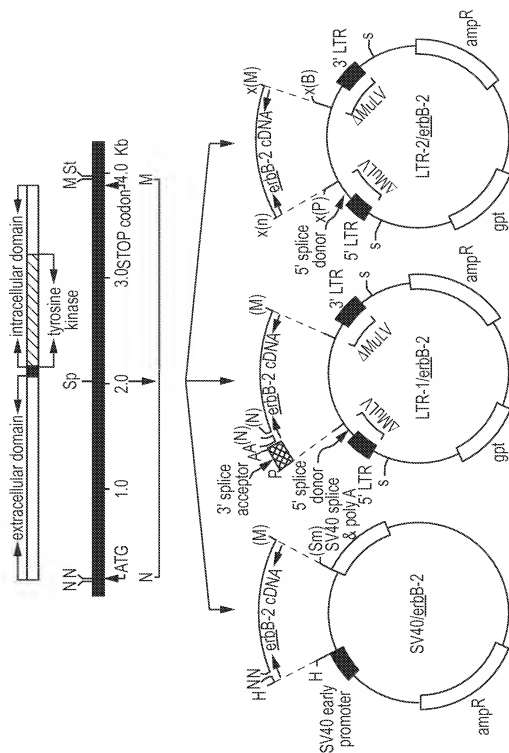


FIG.9

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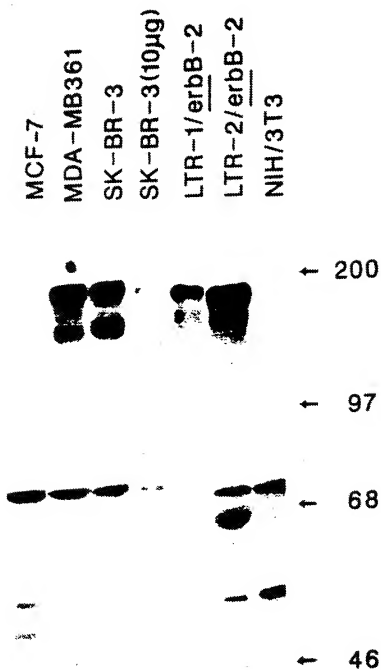


FIG.10